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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.	
10/538,002	06/13/2006	Roberto A Macina	DEX-0547	8194	
	2590 01/14/2008		EXAM	INER	
LICATA & TYRRELL P.C. 66 E. MAIN STREET			MARTINELL, JAMES		
MARLTON, NJ 08053			ART UNIT	PAPER NUMBER	
			1634		
			NOTIFICATION DATE	DELIVERY MODE	
			01/14/2008	ELECTRONIC	

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

Notice of the Office communication was sent electronically on above-indicated "Notification Date" to the following e-mail address(es):

poreilly@licataandtyrrell.com

r	Application No.	Applicant(s)			
đ		MACINA ET AL.			
Office Action Summary	10/538,002	Art Unit			
•	Examiner				
The MAILING DATE of this communication app	James Martinell	1634			
Period for Reply		*			
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period w - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION (6(a). In no event, however, may a reply be tirgoing apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	N. nely filed the mailing date of this communication. ED (35 U.S.C. § 133).			
Status					
1) Responsive to communication(s) filed on 05 No.	ovember 2007.				
2a) ☐ This action is FINAL . 2b) ☒ This	action is non-final.				
3) Since this application is in condition for allowar					
closed in accordance with the practice under E	x parte Quayle, 1935 C.D. 11, 4	53 O.G. 213.			
Disposition of Claims					
4)⊠ Claim(s) <u>1-18</u> is/are pending in the application.					
4a) Of the above claim(s) 11-14 is/are withdraw	n from consideration.				
5) Claim(s) is/are allowed.					
6)⊠ Claim(s) <u>1-10 and 15-18</u> is/are rejected.					
7) Claim(s) is/are objected to.	· · · · · · · · · · · · · · · · · · ·				
8) Claim(s) are subject to restriction and/or	r election requirement.				
Application Papers					
9) The specification is objected to by the Examine	r.	•			
10)☐ The drawing(s) filed on is/are: a)☐ acce	epted or b) objected to by the	Examiner.			
Applicant may not request that any objection to the					
Replacement drawing sheet(s) including the correction					
11)☐ The oath or declaration is objected to by the Ex	aminer. Note the attached Office	e Action or form PTO-152.			
Priority under 35 U.S.C. § 119					
12) Acknowledgment is made of a claim for foreign a) All b) Some * c) None of: 1. Certified copies of the priority documents	s have been received.				
 2. Certified copies of the priority documents have been received in Application No 3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). 					
* See the attached detailed Office action for a list of the certified copies not received.					
	•				
Attachment(s)					
 Notice of References Cited (PTO-892) Notice of Draftsperson's Patent Drawing Review (PTO-948) Information Disclosure Statement(s) (PTO/SB/08) Paper No(s)/Mail Date 6/3/05. 	4) Interview Summan Paper No(s)/Mail D 5) Notice of Informal 6) Other:	oate			
		<u></u>			

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Applicant's election with traverse of the requirement for restriction in the reply filed on November 5, 2007 is acknowledged. The traversal is on the ground(s) that "a search of the art relating to an elected nucleic acid sequence should reveal art relating to the protein encoded thereby and antibodies thereto". This is not found persuasive because the searches of the three Groups of inventions are not coextensive. It is noted that applicants did not argue against the selection of a single sequence for examination on the merits.

The requirement is still deemed proper and is therefore made FINAL.

Claims 11-14, 15 (insofar as it is drawn to polypeptide assays) and 16-18 (insofar as they are drawn to kits containing polypeptides (claim 16), methods of treatment using polypeptides (claim 17), and polypeptide vaccines (claim 18)) are withdrawn from further consideration pursuant to 37 CFR 1.142(b), as being drawn to a nonelected invention, there being no allowable generic or linking claim. Applicant timely traversed the restriction (election) requirement in the reply filed on November 5, 2007.

The disclosure is objected to because it contains an embedded hyperlink and/or other form of browser-executable code. Applicant is required to delete the embedded hyperlink and/or other form of browser-executable code. See MPEP §608.01. Embedded hyperlink and/or other form of browser-executable code appear in at least the following locations:

(a) page 195, line 6.

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 1-10 and 15-18 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention. The claims are vague, indefinite, and incomplete.

(a) Claims 1 and 15-18 are vague and indefinite because they claim more than was elected. Claims 1, 15, 16, and 17 are drawn to or require the use of

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more than one selected nucleic acid sequence. Claim 15 (parts (a) (v) and (vi) and part (b) comparing polypeptide amounts) is drawn to a non-elected invention. Claim 16 (part (e)) is drawn to kits containing polypeptides.

Claim 17 (part (e)) is drawn to methods of using polypeptides. Part of claim 18 is drawn to polypeptide vaccines.

- (b) The recitation of "selectively hybridizes" (claims 1, 15, 16, and 17) is vague, indefinite, and incomplete because nucleic acid molecular hybridization is a process in which selective hybridization is dependent upon competing target in the hybridization mixture (*e.g.*, see Kennell (Progr. Nucl. Acid Res. Mol. Biol. 11: 259 (1971)) cited here as of interest). Since the claims give no information about the presence or absence of competing targets, the claims are vague, indefinite, and incomplete. The metes and bounds of the claims are not clear.
- incomplete because nucleic acid molecular hybridization is a process in which selective hybridization is dependent upon competing target in the hybridization mixture (*e.g.*, see Kennell (Progr. Nucl. Acid Res. Mol. Biol. 11: 259 (1971)) cited here as of interest). Since the claim gives no information about the presence or absence of competing targets, the claim is vague, indefinite, and incomplete. The metes and bounds of the claim are not clear.
- (d) Claims 16, 17, and 18 are incomplete because they depend from cancelled claim 12.
- (e) Claims 7 and 10 are vague and indefinite because they are improper hybrid claims. Each of claims 7 and 10 is a method claim, while claim 1, from which each of claims 7 and 10 depends, is a composition claim.

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The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

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Claims 1-6, 8, 9, and 10 are rejected under 35 U.S.C. 102(b) as being clearly anticipated by either one of Stormann et al (U.S. Patent No. 6,077,675) or Drmanac et al (WO 01/75067 A2 (October 11, 2001)). Stormann et al discloses a nucleic acid that shares 3589 nucleotides with SEQ ID NO: 9 of the instant claims (see the alignment below). Thus, the nucleic acids of Stormann et al would hybridize to SEQ ID NO: 9 of the instant claims. Kennell (Progr. Nucl. Acid Res. Mol. Biol. 11: 259 (1971) paragraph bridging pages 260-261) teaches that a heteroduplex of 25-50 base pairs approaches maximal stability, thus, the nucleic acids of Stormann et al are embraced by the claims. In addition, Stormann et al teaches the use of vectors and host cells to produce a polypeptide encoded by the nucleic acid (*e.g.*, see column 11, line 20 through column 12, line 27).

Alignment of Stormann et al SEQ ID NO: 1 with SEQ ID NO: 9

```
RESULT 1
US-08-823-110-2
; Sequence 2, Application US/08823110
; Patent No. 6077675
  GENERAL INFORMATION:
     APPLICANT:
                Stormann, Thomas M.
     APPLICANT: Simin, Rachel T.
    APPLICANT: Hammerland, Lance G.
     APPLICANT: Fuller, Forrest H.
     TITLE OF INVENTION: NOVEL HUMAN METABOTROPIC
     TITLE OF INVENTION: GLUTAMATE RECEPTOR
     NUMBER OF SEQUENCES: 16
    CORRESPONDENCE ADDRESS:
       ADDRESSEE: Lyon & Lyon
       STREET: 633 West.Fifth Street
       STREET: Suite 4700
      CITY: Los Angeles
STATE: California
      COUNTRY: U.S.A.
       ZIP: 90071-2066
    COMPUTER READABLE FORM:
      MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
      MEDIUM TYPE: storage
      COMPUTER: IBM Compatible
       OPERATING SYSTEM: IBM P.C. DOS 5.0
       SOFTWARE: FastSEQ for Windows 2.0
```

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CURRENT APPLICATION DATA:
     APPLICATION NUMBER: US/08/823,110
     FILING DATE: March 24, 1997
     CLASSIFICATION:
   PRIOR APPLICATION DATA:
     APPLICATION NUMBER: 08/604,298
     FILING DATE: February 21, 1996
   ATTORNEY/AGENT INFORMATION:
     NAME: Warburg, Richard J.
     REGISTRATION NUMBER: 32,327
     REFERENCE/DOCKET NUMBER: 224/259
   TELECOMMUNICATION INFORMATION:
     TELEPHONE: (213) 489-1600
     TELEFAX: (213) 955-0440
     TELEX: 67-3510
  INFORMATION FOR SEQ ID NO: 2:
   SEQUENCE CHARACTERISTICS:
     LENGTH: 3833 base pairs
     TYPE: nucleic acid
     STRANDEDNESS: single
     TOPOLOGY: linear
US-08-823-110-2
 Query Match 91.4%; Score 3412.8; DB 3; Length 3833; Best Local Similarity 96.2%; Pred. No. 0;
                          0; Mismatches
 Matches 3589; Conservative
                                       12;
                                           Indels 129; Gaps
         1 GTGTGCTGGAAAGGCGTGCCCCGCTGGGCTTCTTCACCGCGCCCTCTGCGGGGAGCAGG 60
Ov
              145 GTGACATGGAGCTGCGGGCCCCCGGCGGGGCTTCCTCACCGCGCCCTCTGCGGGGAGCAGG 204
Db
        61 GAATAATTCTGCTACAAGGCTGATTTCAAGGACATGAATTGTTGACCTCATCCCAACATC 120
Qу
           205 GAATAATTCTGCTACAAGGCTGATTTCAAGGACATGAATTGTTGACCTCATCCCAACATC 264
Db
        121 AGAACCTCAGATGTTCTAATTTTTGCACCATTCCAGGCAAGTTGATCTTATAAGGAAATA 180
Qу
           265 AGAACCTCAGATGTTCTAATTTTTGCACCATTCCAGGCAAGTTGATCTTATAAGGAAATA 324
Db
        181 AAATTGAACCTTAGGGGTCTGATGGAAATTCACTGTGACATTCAAATCAAGAAAACTTGC 240
0v
           325 AAATTGAACCTTAGGGGTCTGATGGAAATTCACTGTGACATTCAAATCAAGAAAACTTGC 384
Db
        241 TAATGCCCACAGAGCCTTTTCCCCATGGGCCCTGATGGTAGCCTCCAGAAGGTGCAGCCT 300
Qy
           385 TAATGCCCACAGAGCCTTTTCCCCATGGGCCCTGATGGTAGCCTCCAGAAGGTGCAGCCT 444
        301 CAGGTGGTGCCCTTTCTGCTGTTTGCAAGAATAAACTTTGGGTCTTGGATTGCAATACC 360
Qν
           445 CAGGTGGTGCCCTTTCTTCTGTG--GCAAGAATAAACTTTGGGTCTTGGATTGCAATACC 502
Db
        361 ACCTGTGGAGAAAATGGTATGCGAGGGAAAGCGATCAGCCTCTTGCCCTTGTTTCTTCCT 420
Qу
           503 ACCTGTGGAGAAATGGTATGCGAGGGAAAGCGATCAGCCTCTTGCCCTTGTTTCTTCCT 562
Db
        421 CTTGACCGCCAAGTTCTACTGGATCCTCACAATGATGCAAAGAACTCACAGCCAGGAGTA 480
Qy
           563 CTTGACCGCCAAGTTCTACTGGATCCTCACAATGATGCAAAGAACTCACAGCCAGGAGTA 622
Db
        481 TGCCCATTCCATACGGGTGGATGGGGACATTATTTTGGGGGGGTCTCTTCCCTGTCCACGC 540
Qу
           623 TGCCCATTCCATACGGGTGGATGGGGACATTATTTTGGGGGGGTCTCTTCCCTGTCCACGC 682
Db
        541 AAAGGGAGAGAGGGGTGCCTTGTGGGGAGCTGAAGAAGGAAAAGGGGATTCACAGACT 600
0ν
           AAAGGGAGAGAGAGGGGTGCCTTGTGGGGAGCTGAAGAAGGAAAAGGGGATTCACAGACT 742
Db
```

Qy	601	${\tt GGAGGCCATGCTTTATGCAATTGACCAGATTAACAAGGACCCTGATCTCCTTTCCAACAT}$	660
Db	743		802
Qy	661	CACTCTGGGTGTCCGCATCCTCGACACGTGCTCTAGGGACACCTATGCTTTGGAGCAGTC	720
Db	803		862
Qу	721	${\tt TCTAACATTCGTGCAGGCATTAATAGAGAAAGATGCTTCGGATGTGAAGTGTGCTAATGG}$	780
Db	863		922
Qу		AGATCCACCCATTTTCACCAAGCCCGACAAGATTTCTGGCGTCATAGGTGCTGCAGCAAG	
Db	923	AGATCCACCCATTTTCACCAAGCCCGACAAGATTTCTGGCGTCATAGGTGCTGCAGCAAG	982
Qy	841	CTCCGTGTCCATCATGGTTGCTAACATTTTAAGACTTTTTAAGATACCTCAAATCAGCTA	900
Db	983	$\tt CTCCGTGTCCATCATGGTTGCTAACATTTTAAGACTTTTTAAGATACCTCAAATCAGCTA$	1042
Qу	901	TGCATCCACAGCCCCAGAGCTAAGTGATAACACCAGGTATGACTTTTTCTCTCGAGTGGT	960
Db	1043	${\tt TGCATCCACAGCCCCAGAGCTAAGTGATAACACCAGGTATGACTTTTTCTCTCGAGTGGT}$	1102
Qу	961	TCCGCCTGACTCCTACCAAGCCCAAGCCATGGTGGACATCGTGACAGCACTGGGATGGAA	1020
Db	1103		1162
Qy .	1021	$\tt TTATGTTTCGACACTGGCTTCTGAGGGGAACTATGGTGAGAGCGGTGTGGAGGCCTTCAC$	1080
Db	1163		1222
Qy	1081	CCAGATCTCGAGGGAGATTGGTGGTGTTTGCATTGCTCAGTCACAGAAAATCCCACGTGA	1140
Db	1223		1282
Qу	1141	ACCAAGACCTGGAGAATTTGAAAAAATTATCAAACGCCTGCTAGAAACACCTAATGCTCG	1200
Db	1283		1342
Qy	1201	AGCAGTGATTATGTTTGCCAATGAGGATGACATCAGGAGGATATTGGAAGCAGCAAAAAA	1260
Db	1343		1402
Qy	1261	ACTAAACCAAAGTGGGCATTTTCTCTGGATTGGCTCAGATAGTTGGGGATCCAAAATAGC	1320
Db	1403		1462
Qy		ACCTGTCTATCAGCAAGAGGAGATTGCAGAAGGGGCTGTGACAATTTTGCCCAAACGAGC	
Db			
		ATCAATTGATGGATTTGATCGATACTTTAGAAGCCGAACTCTTGCCAATAATCGAAGAAA	
Qy			
Db			
Qу		TGTGTGGTTTGCAGAATTCTGGGAGGAGAATTTTGGCTGCAAGTTAGGATCACATGGGAA	
Db		TGTGTGGTTTGCAGAATTCTGGGAGGAGAATTTTGGCTGCAAGTTAGGATCACATGGGAA	
Qу		AAGGAACAGTCATATAAAGAAATGCACAGGGCTGGAGCGAATTGCTCGGGATTCATCTTA	
Db	1643	AAGGAACAGTCATATAAAGAAATGCACAGGGCTGGAGCGAATTGCTCGGGATTCATCTTA	1702

Qy	1561	TGAACAGGAAGGTCCAATTTGTAATTGATGCTGTATATTCCATGGCTTACGCCCT	1620
Db	1703	TGAACAGGAAGGAAAGGTCCAATTTGTAATTGATGCTGTATATTCCATGGCTTACGCCCT	1762
Qy	1621	GCACAATATGCACAAAGATCTCTGCCCTGGATACATTGGCCTTTGTCCACGAATGAGTAC	1680
Db	1763		1822
Qу	1681	CATTGATGGGAAAGAGCTACTTGGTTATATTCGGGCTGTAAATTTTAATGGTTGCCGAAG	1740
Db	1823	CATTGATGGGAAAGAGCTACTTGGTTATATTCGGGCTGTAAATTTTAAT	1871
Qy	1741	AGGGATCCAGATGTCTCTACCCTGGCCAACTCTTTTTACTCCTTCATTTTCCAGTAGTTG	1800
Db			
Qy		GGCAGTGCTGGCACTCCTGTCACTTTTAATGAAAACGGAGATGCTCCTGGACGTTATGAT	
w1 Db		GGAGTGCTGGCACTCCTGTCACTTTTAATGAAAACGGAGATGCTCCTGGACGTTATGAT	
_		ATCTTCCAGTATCAAATAACCAACAAAAGCACAGAGTACAAAGTCATCGGCCACTGGACC	
Qy Db			
_		AATCAGCTTCATCTAAAAGTGGAAGACATGCAGTGGGCTCATAGAGAACATACTCACCCG	
Qy Db		AATCAGCTTCATCTAAAAGTGGAAGACATGCAGTGGGCTCATAGAGMACATACTCACCCG AATCAGCTTCATCTAAAAGTGGAAGACATGCAGTGGGCTCATAGAGAACATACTCACCCG	
		GCGTCTGTCTGCAGCCTGCCGTGTAAGCCAGGGGAGGAAGAAAACGGTGAAAGGGGTC	
Qy		GCGTCTGTCTGCAGCCTGCCGTGTAAGCCAGGGGAAGAAAACGGTGAAAGGGGTC 	
Db		CCTTGCTGCTGCAGCCTGTGAACGCTGTGAAGGTTACAACTACCAGGTGGATGAGCTGTCC	
Qу		CCTTGCTGCTGCACCTGTGAACGCTGTGAAGGTTACAACTACCAGGTGGATGAGCTGTCC	
Db		•	
Qy 		TGTGAACTTTGCCCTCTGGATCAGAGACCCAACATGAACCGCACAGGCTGCCAGCTTATC	
Db		TGTGAACTTTGCCCTCTGGATCAGAGACCCAACATGAACCGCACAGGCTGCCAGCTTATC	
Qy		CCCATCATCAAATTGGAGTGGCATTCTCCCTGGGCTGTGTGTTTGTT	
Db		CCCATCATCAAATTGGAGTGGCATTCTCCCTGGGCTGTGGTGCCTGTGTTTGTT	
Qу		TTGGGAATCATCGCCACCACCTTTGTGATCGTGACCTTTGTCCGCTATAATGACACACCT	
Db		TTGGGAATCATCGCCACCACCTTTGTGATCGTGACCTTTGTCCGCTATAATGACACACCT	
Qу		ATCGTGAGGGCTTCAGGACGCGAACTTAGTTACGTGCTCCTAACGGGGATTTTTCTCTGT	•
Db		ATCGTGAGGGCTTCAGGACGCGAACTTAGTTACGTGCTCCTAACGGGGATTTTTCTCTGT	
Qy		TATTCAATCACGTTTTTAATGATTGCAGCACCAGATACAATCATATGCTCCTTCCGACGG	
Db		TATTCAATCACGTTTTTAATGATTGCAGCACCAGATACAATCATATGCTCCTTCCGACGG	
Qу		GTCTTCCTAGGACTTGGCATGTTTTCAGCTATGCAGCCCTTCTGACCAAAACAAAC	
Db	2472	2 GTCTTCCTAGGACTTGGCATGTGTTTCAGCTATGCAGCCCTTCTGACCAAAACAAAC	2531
Qу	2461	ATCCACCGAATATTTGAGCAGGGGAAGAAATCTGTCACAGCGCCCAAGTTCATTAGTCCA	2520
Db	2532	ATCCACCGAATATTTGAGCAGGGGAAGAAATCTGTCACAGCGCCCAAGTTCATTAGTCCA	2591
Ov	2521	GCATCTCAGCTGGTGATCACCTTCAGCCTCATCTCCGTCCAGCTCCTTGGAGTGTTTGTC	2580

		1,	
Db	2592	GCATCTCAGCTGGTGATCACCTTCAGCCTCATCTCCGTCCAGCTCCTTGGAGTGTTTGTC	2651
Qу	2581	TGGTTTGTTGTGGATCCCCCCACATCATCATTGACTATGGAGAGCAGCGGACACTAGAT	2640
Db	2652	TGGTTTGTTGTGGATCCCCCCCACATCATCATTGACTATGGAGAGCAGCGGACACTAGAT	2711
Qу	2641	CCAGAGAAGGCCAGGGGAGTGCTCAAGTGTGACATTTCTGATCTCTCACTCA	2700
Db	2712	CCAGAGAAGGCCAGGGGAGTGCTCAAGTGTGACATTTCTGATCTCTCACTCA	2771
Qу	2701	CTTGGATACAGTATCCTCTTGATGGTCACTTGTACTGTTTATGCCATTAAAACGAGAGGT	2760
Db	2772	CTTGGATACAGTATCCTCTTGATGGTCACTTGTACTGTTTATGCCATTAAAACGAGAGGT	2831
Qy	2761	GTCCCAGAGACTTTCAATGAAGCCAAACCTATTGGATTTACCATGTATACCACCTGCATC	2820
Db .	2832	GTCCCAGAGACTTTCAATGAAGCCAAACCTATTGGATTTACCATGTATACCACCTGCATC	2891
Qy	2821	ATTTGGTTAGCTTCATCCCCATCTTTTTTGGTACAGCCCAGTCAGCAGAAAAGATGTAC	2880
Db	2892	ATTTGGTTAGCTTTCATCCCCATCTTTTTTGGTACAGCCCAGTCAGCAGAAAAGATGTAC	2951
Qy	2881	ATCCAGACAACACTTACTGTCTCCATGAGTTTAAGTGCTTCAGTATCTCTGGGCATG	2940
Db	2952	ATCCAGACAACACTTACTGTCTCCATGAGTTTAAGTGCTTCAGTATCTCTGGGCATG	3011
Qy	2941	CTCTATATGCCCAAGGTTTATATATATTTTTCATCCAGAACAGAATGTTCAAAAACGC	3000
Db	3012	CTCTATATGCCCAAGGTTTATATTATAATTTTTCATCCAGAACAGAATGTTCAAAAACGC	3071
Qy	3001	AAGAGGAGCTTCAAGGCTGTGGTGACAGCTGCCACCATGCAAAGCAAACTGATCCAAAAA	3060
Db	3072	AAGAGGAGCTTCAAGGCTGTGGTGACAGCTGCCACCATGCAAAGCAAACTGATCCAAAAA	3131
Qу	3061	GGAAATGACAGACCAAATGGCGAGGTGAAAAGTGAACTCTGTGAGAGTCTTGAAACCAAC	3120
Db	3132	GGAAATGACAGACCAAATGGCGAGGTGAAAAGTGAACTCTGTGAGAGTCTTGAAACCAAC	3191
Qy	3121	ACTTC	3125
Db	3192	AGTAAGTCATCTGTAGAGTTTCCGATGGTCAAGAGCGGGAGCACTTCCTAATAGATCTTC	3251
Qу	3126	CTCTACCAAGACAACATATATCAGTTACAGCAATCATTCAATCTGAAACAGGGAAATGGC	3185
Db	3252	CTCTACCAAGACATATATCAGTTACAGCAATCATTCAATCTGAAACAGGGAAATGGC	3311
Qу	3186	ACAATCTGAAGAGATGTGGTATATGATCTTAAATGATGAACATGAGACCGCAAAAATTCA	3245
Db	3312	ACAATCTGAAGAGACGTGGTATATGATCTTAAATGATGAACATGAGACCGCAAAAATTCA	3371
Qу	3246	CTCCTGGAGATCTCCGTAGACTACAATCAATCAATCAATAGTCAGTC	3305
Db	3372	CTCCTGGAGATCTCCGTAGACTACAATCAATCAATAGTCAGTC	3431
Qу	3306	AAAATTAGCCATGAGCCAAAAGTATCAATAAACGGGGAGTGAAGAAACCCGTTTTATACA	3365
Db	3432	AAAATTAGCCATGAGCCAAAAGTATCAATAAACGGGGAGTGAAGAAACCCGTTTTATACA	3491
Qy	3366	ATAAAACCAATGAGTGTCAAGCTAAAGTATTGCTTATTCATGAGCAGTTAAAACAAATCA	3425
Db	3492	ATAAAACCAATGAGTGTCAAGCTAAAGTATTGCTTATTCATGAGCAGTTAAAACAAATCA	3551
Qy	3426	CAAAAGGAAAACTAATGTTAGCTCGTGAAAAAAAATGCTGTTGAAATAAAT	3485
Db	3552	CAAAAGGAAAACTAATGTTAGCTCGTG-AAAAAAATGCTGTTGAAATAAATAATGTCTGA	3610

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Qу
     3486 TGTTATTCTTGTATTTTTCTGTGATTGTGAGAACTCCCGTTCCTGTCCCACATTGTTTAA 3545
         3611 TGTTATTCTTGTATTTTTCTGTGATTGTGAGAACTCCCGTTCCTGTCCCACATTGTTTAA 3670
Db
Qу
     3546 CTTGTATAAGACAATGAGTCTGTTTCTTGTAATGGCTGACCAGATTGAAGCCCTGGGTTG 3605
         Db
     3671 CTTGTATAAGACAATGAGTCTGTTTCTTGTAATGGCTGACCAGATTGAAGCCCTGGGTTG 3730
     Qy
         Db
     3731 TGCTAAAAATAAATGCAATGATTGATGCATGCAATTTTTTATACAAATAATTTATTCTA 3790
     3666 ATAATAAAGG 3675
Qу
        3791 ATAATAAAGG 3800
Db
```

Claims 1-6, 8-10, 16, and 18 are rejected under 35 U.S.C. 102(b) as being clearly anticipated by Drmanac et al (WO 01/75067 A2 (October 11, 2001)). Drmanac et al discloses a nucleic acid (SEQ ID NO: 29442) that shares 3351 nucleotides with SEQ ID NO: 9 of the instant claims (see the alignment below). Thus, the nucleic acids of Drmanac et al would hybridize to SEQ ID NO: 9 of the instant claims. Kennell (Progr. Nucl. Acid Res. Mol. Biol. 11: 259 (1971) paragraph bridging pages 260-261) teaches that a heteroduplex of 25-50 base pairs approaches maximal stability, thus, the nucleic acids of Drmanac et al are embraced by the claims. In addition, Drmanac et al teaches the use of vectors and host cells to produce a polypeptide encoded by the nucleic acid (*e.g.*, see page 19, line 7 through page 20, line 24), the use of polynucleotide vaccines (*e.g.*, page 20, lines 17-24), and the inclusion of polynucleotides in kits (*e.g.*, page 87, line 20 through page 89, line 5).

Alignment of Drmanac et al SEQ ID NO: 29442 with SEQ ID NO: 9

```
RESULT 3
AAS93638
ID AAS93638 standard; cDNA; 3940 BP.
XX
AC AAS93638;
XX
DT 13-FEB-2002 (first entry)
XX
DE DNA encoding novel human diagnostic protein #29442.
```

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KW

```
Human; chromosome mapping; gene mapping; gene therapy; forensic;
KW
    food supplement; medical imaging; diagnostic; genetic disorder; ss.
XX
os
    Homo sapiens.
XX
PN
    W0200175067-A2.
XX
PD
    11-OCT-2001.
XX
    30-MAR-2001; 2001WO-US008631.
PF
PR
    31-MAR-2000; 2000US-00540217.
PR
    23-AUG-2000; 2000US-00649167.
XX
PA
     (HYSE-) HYSEQ INC.
XX
PΙ
    Drmanac RT, Liu C, Tang YT;
XX
DR
    WPI; 2001-639362/73.
DR
    P-PSDB; ABG29451.
XX
    New isolated polynucleotide and encoded polypeptides, useful in
    diagnostics, forensics, gene mapping, identification of mutations
РΤ
PT
    responsible for genetic disorders or other traits and to assess
PT
    biodiversity.
    Claim 1; SEQ ID NO 29442; 103pp; English.
PS
XX
    The invention relates to isolated polynucleotide (I) and polypeptide (II)
CC
CC
    sequences. (I) is useful as hybridisation probes, polymerase chain
CC
    reaction (PCR) primers, oligomers, and for chromosome and gene mapping,
    and in recombinant production of (II). The polynucleotides are also used
CC
CC
    in diagnostics as expressed sequence tags for identifying expressed
CC
    genes. (I) is useful in gene therapy techniques to restore normal
CC
    activity of (II) or to treat disease states involving (II). (II) is
CC
    useful for generating antibodies against it, detecting or quantitating a
CC
    polypeptide in tissue, as molecular weight markers and as a food
    supplement. (II) and its binding partners are useful in medical imaging
CC
CC
    of sites expressing (II). (I) and (II) are useful for treating disorders
    involving aberrant protein expression or biological activity. The
CC
    polypeptide and polynucleotide sequences have applications in
CC
CC
    diagnostics, forensics, gene mapping, identification of mutations
CC
    responsible for genetic disorders or other traits to assess biodiversity
CC
    and to produce other types of data and products dependent on DNA and
    amino acid sequences. AAS64197-AAS94564 represent novel human diagnostic
CC
    coding sequences of the invention. Note: The sequence data for this
CC
    patent did not appear in the printed specification, but was obtained in
CC
CC
    electronic format directly from WIPO at
CC
    ftp.wipo.int/pub/published_pct_sequences
XX
    Sequence 3940 BP; 1071 A; 921 C; 952 G; 996 T; 0 U; 0 Other;
SO
                        89.2%; Score 3331; DB 5; Length 3940;
 Query Match
                        99.5%; Pred. No. 0;
 Best Local Similarity
                               0; Mismatches
                                                                         1:
 Matches 3351; Conservative
                                              15: Indels
                                                                 Gaps
         309 GCCCTTTCTGCTGTTGCAAGAATAAACTTTGGGTCTTGGATTGCAATACCACCTGTGG 368
Qу
                  11111
                        566 GCGGCTTCCGGGATGAGTCGGGAATAAACTTTGGGTCTTGGATTGCAATACCACCTGTGG 625
Db
         369 AGAAATGGTATGCGAGGGAAAGCGATCAGCCTCTTGCCCTTGTTTCTTCCTCTTGACCG 428
Qy
             626 AGAAATGGTATGCGAGGGAAAGCGATCAGCCTCTTGCCCTTGTTTCTTCCTCTTGACCG 685
Db
         429 CCAAGTTCTACTGGATCCTCACAATGATGCAAAGAACTCACAGCCAGGAGTATGCCCATT 488
Qy
             686 CCAAGTTCTACTGGATCCTCACAATGATGCAAAGAACTCACAGCCAGGAGTATGCCCATT 745
Db
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Qу	489	$\verb CCATACGGGTGGATGGGGACATTATTTTGGGGGGTCTCTTCCCTGTCCACGCAAAGGGAG \\$	548
Db	746		805
Qy	549	${\tt AGAGAGGGGTGCCTTGTGGGGAGCTGAAGAAGGAAAAGGGGATTCACAGACTGGAGGCCA}$	608
Db	806		865
Qу	609	$\tt TGCTTTATGCAATTGACCAGATTAACAAGGACCCTGATCTCCTTTCCAACATCACTCTGG$	668
Db	866	TGCTTTATGCAATTGACCAGATTAACAAGGACCCTGATCTCCTTTCCAACATCACTCTGG	925
Qу	669	GTGTCCGCATCCTCGACACGTGCTCTAGGGACACCTATGCTTTGGAGCAGTCTCTAACAT	728
Db	926		985
Qу	729	TCGTGCAGGCATTAATAGAGAAAGATGCTTCGGATGTGAAGTGTGCTAATGGAGATCCAC	788
Db	986	TCGTGCAGGCATTAATAGAGAAAGATGCTTCGGATGTGAAGTGTGCTAATGGAGATCCAC	1045
Qy	789	$\verb CCATTTTCACCAAGCCCGACAAGATTTCTGGCGTCATAGGTGCTGCAGCAAGCTCCGTGT \\$	848
Db	1046	CCATTTCACCAAGCCCGACAAGATTTCTGGCGTCATAGGTGCTGCAGCAAGCTCCGTGT	1105
Qу	849	${\tt CCATCATGGTTGCTAACATTTTAAGACTTTTTAAGATACCTCAAATCAGCTATGCATCCA}$	908
Db	1106		1165
Qу	909	${\tt CAGCCCCAGAGCTAAGTGATAACACCAGGTATGACTTTTTCTCTCGAGTGGTTCCGCCTG}$	968
Db	1166		1225
Qу	969	${\tt ACTCCTACCAAGCCCAAGCCATGGTGGACATCGTGACAGCACTGGGATGGAATTATGTTT}$	1028
Db	1226	ACTCCTACCAAGCCCAAGCCATGGTGGACATCGTGACAGCACTGGGATGGAATTATGTTT	1285
Qу	1029	$\tt CGACACTGGCTTCTGAGGGGAACTATGGTGAGAGGGGTGTGGAGGCCTTCACCCAGATCT$	1088
Db	1286	CGACACTGGCTTCTGAGGGGAACTATGGTGAGACGGTGTGGAGGCCTTCACCCAGATCT	1345
Qy	1089	${\tt CGAGGGAGATTGGTGTTTTGCATTGCTCAGTCACAGAAAATCCCACGTGAACCAAGAC}$	1148
Db	1346	CGAGGGAGATTGGTGTTTTGCATTGCCCAGTCACAGAAAATCCCACGTGAACCAAGAC	1405
Qy	1149	CTGGAGAATTTGAAAAAATTATCAAACGCCTGCTAGAAACACCTAATGCTCGAGCAGTGA	1208
Db	1406	CTGGAGAATTTGAAAAATTATCAAACGCCTGCTAGAAACACCTAATGCTCGAGCAGTGA	1465
Qy	1209	$\verb"TTATGTTTGCCAATGAGGATGACATCAGGAGGATATTGGAAGCAGCAAAAAAAA$	1268
Db	1466		1525
Qy	1269	${\tt AAAGTGGGCATTTTCTCTGGATTGGCTCAGATAGTTGGGGATCCAAAATAGCACCTGTCT}$	1328
Db	1526		1585
Qy	1329	$\dot{\text{ATCAGCAAGAGGAGATTGCAGAAGGGGCTGTGACAATTTTGCCCAAACGAGCATCAATTG}$	1388
Db	1586		1645
Qy	1389	${\tt ATGGATTTGATCGATACTTTAGAAGCCGAACTCTTGCCAATAATCGAAGAAATGTGTGGGT}$	1448
Db	1646		1705
Qу	1449	${\tt TTGCAGAATTCTGGGAGGAGAATTTTGGCTGCAAGTTAGGATCACATGGGAAAAGGAACA}$	1508
Db	1706		1765

Qу	1509	GTCATATAAAGAAATGCACAGGGCTGGAGCGAATTGCTCGGGÄTTCATCTTATGAACAGG	1568
Db	1766	GTCATATAAAGAAATGCACAGGGCTGGAGCGAATTGCTCGGGATTCATCTTATGAACAGG	1825
Qy	1569	AAGGAAAGGTCCAATTTGTAATTGATGCTGTATATTCCATGGCTTACGCCCTGCACAATA	1628
Db	1826	AAGGAAAGGTCCAATTTGŤAATTGATGCTGTATATTCCATGGCTTACGCCCTGCACAATA	1885
Qy	1629	TGCACAAAGATCTCTGCCCTGGATACATTGGCCTTTGTCCACGAATGAGTACCATTGATG	1688
Db	1886		1945
Qy	1689	GGAAAGAGCTACTTGGTTATATTCGGGCTGTAAATTTTAATGGTTGCCGAAGAGGGATCC	1748
Db	1946		2005
Qу	.1749	${\tt AGATGTCTCTACCCTGGCCAACTCTTTTTACTCCTTCATTTTCCAGTAGTTGGGCAGTGC}$	1808
Db	2006		2065
Qy	1809	${\tt TGGCACTCCTGTCACTTTTAATGAAAACGGAGATGCTCCTGGACGTTATGATATCTTCCA}$	1868
Db	2066		2125
Qу	1869	GTATCAAATAACCAACAAAAGCACAGAGTACAAAGTCATCGGCCACTGGACCAATCAGCT	1928
Db	2126	GTATCAAATAACCAACAAAAGCACAGAGTACAAAGTCATCGGCCACTGGACCAATCAGCT	2185
Qу	1929	TCATCTAAAAGTGGAAGACATGCAGTGGGCTCATAGAGAACATACTCACCCGGCGTCTGT	1988
Db	2186	TCATCTAAAAGTGGAAGACATGCAGTGGGCTCATAGAGAACATACTCACCCGGCGTCTGT	2245
Qу	1989	CTGCAGCCTGCCGTGTAAGCCAGGGGAGGAAGAAAACGGTGAAAGGGGTCCCTTGCTG	2048
Db	2246	$\tt CTGCAGCCTGCCGTGTAAGCCAGGGGAGGAGGAGAAAACGGTGAAAGGGGTCCCTTGCTG$	2305
Qу	2049	CTGGCACTGTGAACGCTGTGAAGGTTACAACTACCAGGTGGATGAGCTGTCCTGTGAACT	2108
Db	2306	CTGGCACTGTGAACGCTGTGAAGGTTACAACTACCAGGTGGATGAGCTGTCCTGTGAACT	2365
Qу	2109	TTGCCCTCTGGATCAGAGACCCAACATGAACCGCACAGGCTGCCAGCTTATCCCCATCAT	2168
Db	2366	$\tt TTGCCCTCTGGATCAGAGACCCAACATGAACCGCACAGGCTGCCAGCTTATCCCCATCAT$	2425
Qу	2169	CAAATTGGAGTGCCATTCTCCCTGGGCTGTGGTGCCTGTGTTTGTT	2228
Db		${\tt CAAATTGGAGTGGCATTCTCCCTGGGCTGTGTGGTGTTTGTT$	
Qy	2229	CATCGCCACCACCTTTGTGATCGTGACCTTTGTCCGCTATAATGACACACCTATCGTGAG	2288
Db		${\tt CATCGCCACCACCTTTGTGATCGTGACCTTTGTCCGCTATAATGACACACCTATCGTGAG}$	
Qу		GGCTTCAGGACGCGAACTTAGTTACGTGCTCCTAACGGGGATTTTTCTCTGTTATTCAAT	
Db		${\tt GGCTTCAGGACGCGAACTTAGTTACGTGCTCCTAACGGGGATTTTTCTCTGTTATTCAAT}$	
Qу		CACGTTTTTAATGATTGCAGCACCAGATACAATCATATGCTCCTTCCGACGGGTCTTCCT	
Db		CACGTTTTTAATGATTGCAGCACCAGATACAATCATATGCTCCTTCCGACGGGTCTTCCT	
Qy		AGGACTTGGCATGTGTTTCAGCTATGCAGCCCTTCTGACCAAAACAAAC	
Db		AGGACTTGGCATGTTTTCAGCTATGCAGCCCTTCTGACCAAAACAAAC	
Qу	2469	AATATTTGAGCAGGGGAAGAAATCTGTCACAGCGCCCAAGTTCATTAGTCCAGCATCTCA	2528

Db	2726	${\tt AATATTTGAGCAGGGGAAGAAATCTGTCACAGCGCCCAAGTTCATTAGTCCAGCATCTCA}$	2785
Qy	2529	${\tt GCTGGTGATCACCTTCAGCCTCATCTCCGTCCAGCTCCTTGGAGTGTTTGTCTGGTTTGT}$	2588
Db	2786	GCTGGTGATCACCTTCAGCCTCATCTCCGTCCAGCTCCTTGGAGTGTTTGTCTGGTTTGT .	2845
Qy .	2589	TGTGGATCCCCCCACATCATCATTGACTATGGAGAGCAGCGGACACTAGATCCAGAGAA	2648
Db	2846	TGTGGATCCCCCCACATCATCATTGACTATGGAGAGCAGCGGACACTAGATCCAGAGAA	2905
Qу	2649	GGCCAGGGGAGTGCTCAAGTGTGACATTTCTGATCTCTCACTCA	2708
Db	2906	GGCCAGGGGAGTGCTCAAGTGTGACATTTCTGATCTCTCACTCA	2965
Qу	2709	CAGTATCCTCTTGATGGTCACTTGTACTGTTTATGCCATTAAAACGAGAGGTGTCCCAGA	2768
Db	2966	CAGTATCCTCTTGATGGTCACTTGTACTGTTTATGCCAATAAAACGAGAGGTGTCCCAGA	3025
Qy .	2769	GACTTTCAATGAAGCCAAACCTATTGGATTTACCATGTATACCACCTGCATCATTTGGTT	2828
Db	3026	GACTTTCAATGAAGCCAAACCTATTGGATTTACCATGTATACCACCTGCATCATTTGGTT	3085
Qy	2829	AGCTTTCATCCCCATCTTTTTTGGTACAGCCCAGTCAGCAGAAAAGATGTACATCCAGAC	2888
Db	3086	AGCTTTCATCCCCATCTTTTTTGGTACAGCCCAGTCAGCAGAAAAGATGTACATCCAGAC	3145
Qy	2889	AACAACACTTACTGTCTCCATGAGTTTAAGTGCTTCAGTATCTCTGGGCATGCTCTATAT	2948
Db	3146	AACAACACTTACTGTCTCCATGAGTTTAAGTGCTTCAGTATCTCTGGGCATGCTCTATAT	3205
Qу	2949	GCCCAAGGTTTATATTATAATTTTTCATCCAGAACAGAA	3008
Db	3206	GCCCAAGGTTTATATTATAATTTTTCATCCAGAACAGAA	3265
Qу	3009	CTTCAAGGCTGTGGCAGCTGCCACCATGCAAAACTGATCCAAAAAGGAAATGA	3068
Db	3266	CTTCAAGGCTGTGGTGACAGCTGCCACCATGCAAAGCAAACTGATCCAAAAAGGAAATGA	3325
Qу	3069	CAGACCAAATGGCGAGGTGAAAAGTGAACTCTGTGAGAGTCTTGAAACCAACACTTCCTC	3128
Db	3326	${\tt CAGACCAAATGGCGAGGTGAAAAGTGAACTCTGTGAGAGTCTTGAAACCAACACTTCCTC}$	3385
Qу	3129	TACCAAGACAACATATATCAGTTACAGCAATCATTCAATCTGAAACAGGGAAATGGCACA	3188
Db	3386	TACCAAGACAACATATATCAGTTACAGCAATCATTCAATCTGAAACAGGGAAATGGCACA	3445
Qу	3189	ATCTGAAGAGATGTGGTATATGATCTTAAATGATGAACATGAGACCGCAAAAATTCACTC	3248
Db	3446	ATCTGAAGAGACGTGGTATATGATCTTAAATGATGAACATGAGACCGCAAAAATTCACTC	3505
Qу		CTGGAGATCTCCGTAGACTACAATCAATCAATCAATAGTCAGTC	
Db	3506	CTGGAGATCTCCGTAGACTACAATCAATCAATCAATAGTCAGTC	3565
Qу		ATTAGCCATGAGCCAAAAGTATCAATAAACGGGGAGTGAAGAAACCCGTTTTATACAATA	
Db		ATTAGCCATGAGCCAAAAGTATCAATAAACGGGGAGTGAAGAAACCCGTTTTATACAATA	
Qу		AAACCAATGAGTGTCAAGCTAAAGTATTGCTTATTCATGAGCAGTTAAAACAAATCACAA	
Db		AAACCAATGAGTGTCAAGCTAAAGTATTGCTTATTCATGAGCAGTTAAAACAAATCACAA	
Qу		AAGGAAAACTAATGTTAGCTCGTGAAAAAAAATGCTGTTGAAATAAAT	
Db	3686	AAGGAAAACTAATGTTAGCTCGTG-AAAAAAATGCTGTTGAAATAAATAATGTCTGATGT	3744

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Qy
     3489 TATTCTTGTATTTTTCTGTGATTGTGAGAACTCCCGTTCCTGTCCCACATTGTTTAACTT 3548
        3745 TATTCTTGTATTTTCTGTGATTGTGAGAACTCCCGTTCCTGTCCCACATTGTTTAACTT 3804
Db
     3549 GTATAAGACAATGAGTCTGTTTCTTGTAATGGCTGACCAGATTGAAGCCCTGGGTTGTGC 3608
Qу
        3805 GTATAAGACAATGAGTCTGTTTCTTGTAATGGCTGACCAGATTGAAGCCCTGGGTTGTGC 3864
Db
     3609 TAAAAATAAATGCAATGATTGATGCATGCAATTTTTTATACAAATAATTTATTCTAATA 3668
Qу
         3669 ATAAAGG 3675
Qу
        111111
Db
     3925 ATAAAGG 3931
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Any inquiry concerning this communication or earlier communications from the examiner should be directed to James Martinell whose telephone number is (571) 272-0719.

The examiner works a flexible schedule and can be reached by phone and voice mail.

Alternatively, a request for a return telephone call may be e-mailed to james.martinell@uspto.gov. Since e-mail communications may not be secure, it is suggested that information in such requests be limited to name, phone number, and the best time to return the call.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Ram Shukla, can be reached on (571) 272-0735.

OFFICIAL FAX NUMBER

The fax phone number for the organization where this application or proceeding is assigned is (571) 273-8300. Any Official Communication to the USPTO should be faxed to this number.

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Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to (571) 272-0547.

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James Martinell, Ph.D. Primary Examiner

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